

**IN THE SPECIFICATION**

**Please re-write the paragraph at page 3, lines 1-13 to read as follows:**

As a result of study for the requirement, PTD (Protein Transduction Domain) was designed. Among various PTDs, the transcription factor, Tat, of Human Immunodeficiency Virus-I, HIV-1 has been studied well. This protein can pass through the cell membrane more effectively when it is organized by part of the amino acids distributed through 47 to 57 (~~YGRKKRRQRRR~~), where positively charged amino acids are distributed, than when it is in a complete form consisting of 87 amino acids [Fawell S. et al. Proc. Natl. Acad. Sci. USA 91, 664-668 (1994)]. Like this, amino acids 267 to 300 of VP22 protein of Herpes Simplex Virus type 1 [Elliott G. et al. Cell, 88, 223-233 (1997)], amino acids 84 to 92 of UL-56 protein of HSV-2 (GeneBank code: DI047[gi:221784]), and amino acids 339 to 355 of ANTP (Antennapedia) protein of Drosophila [Schwarze S.R. et al. Trends Pharmacol Sci. 21, 45-48 (2000)] are examples of other PTDs. Further, artificial peptides comprising positively charged amino acids also showed effects [Laus R. et al. Nature Biotech 18, 1269-1272(2000)].